

*RE-RUN*

## STIC Biotechnology Systems Branch

### RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/473,619C  
Source: IFW16  
Date Processed by STIC: 3-30-05

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



IFW16

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/473,619C

DATE: 03/30/2005  
TIME: 13:48:52

Input Set : A:\CEPH1447.ST25.txt  
Output Set: N:\CRF4\03302005\I473619C.raw

```

3 <110> APPLICANT: Siman, Robert
4     Bozyczko-Coyne, Donna
5     Meyer, Sheryl L.
6     Venkatraman Bhat, Ratan
8 <120> TITLE OF INVENTION: Methods For Detecting Cell Apoptosis
10 <130> FILE REFERENCE: CEPH-1447
12 <140> CURRENT APPLICATION NUMBER: 09/473,619C
13 <141> CURRENT FILING DATE: 1999-12-29
15 <150> PRIOR APPLICATION NUMBER: 60/030,961
16 <151> PRIOR FILING DATE: 1996-11-15
18 <150> PRIOR APPLICATION NUMBER: 08/967,625
19 <151> PRIOR FILING DATE: 1997-11-12
21 <160> NUMBER OF SEQ ID NOS: 2
23 <170> SOFTWARE: PatentIn version 3.0
25 <210> SEQ ID NO: 1
26 <211> LENGTH: 5
27 <212> TYPE: PRT
28 <213> ORGANISM: Artificial Sequence
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W--> 30 <223> OTHER INFORMATION:
W--> 30 <400> 1
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33 1           5
35 <210> SEQ ID NO: 2
36 <211> LENGTH: 7
37 <212> TYPE: PRT
38 <213> ORGANISM: Artificial Sequence
W--> 40 <220> FEATURE:
W--> 40 <223> OTHER INFORMATION:
W--> 40 <400> 2
42 Cys Lys Gly Asp Glu Val Asp
43 1           5

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Or as Not Comply  
Commented Sequence Needed

pls explain  
Source of  
genetic material. (pg.1)

pls insert these

(2) mandatory Numeric identifiers.

Same error

pls insert → mandatory,

<220>-<223>

must be inserted  
whenever <213> is  
Artificial or Unknown



Please See error explanation on page 2.

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/473,619C

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Input Set : A:\CEPH1447.ST25.txt  
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Use of <220> Feature(NEW RULES): Error Explanation ↴  
Sequence(s) are missing the <220> Feature and associated headings.  
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence"  
or "Unknown". Please explain source of genetic material in <220> to <223>  
section (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp.29631-32)  
(Sec.1.823 of new Rules)

Seq#:1,2

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/473,619C

DATE: 03/30/2005  
TIME: 13:48:53

Input Set : A:\CEPH1447.ST25.txt  
Output Set: N:\CRF4\03302005\I473619C.raw

L:30 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:1, <213>

ORGANISM:Artificial Sequence

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ORGANISM:Artificial Sequence

L:30 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:30

L:40 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ#:2, <213>

ORGANISM:Artificial Sequence

L:40 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:2, <213>

ORGANISM:Artificial Sequence

L:40 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:2,Line#:40